## WHAT IS CLAIMED IS:

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- 1. A method of increasing the production of a biologically active compound in a cell wherein the biologically active compound is derived at least in part from methylmalonyl-CoA, the method comprising the step of inhibiting the activity of methylmalonyl-CoA mutase.
- 2. The method of claim 1 wherein the biologically active compound is an immunosuppressant.

3. The method of claim 2 wherein the immunosuppessant is rapamycin, FK520, or ascomycin.

- 4. The method of claim 1 wherein the biologically active compound is an antifungal agent.
  - 5. The method of claim 4 wherein the antifungal agent is rapamycin, candicidin or soraphen.
- 20 6. The method of claim 1 wherein the biologically active compound is an antiparasitic agent.
  - 7. The method of claim 6 wherein the antiparasitic agent is avermectin.
- 25 8. The method of claim 1 wherein the biologically active compound is an antibiotic.
  - 9. The method of claim 8 wherein the antibiotic is a polyketide antibiotic.
- 30 10. The method of claim 9 wherein the polyketide antibiotic is a macrolide polyketide antibiotic.

11. The method of claim 10 wherein the macrolide polyketide antibiotic is erythromycin, tylosin, niddamycin, spiramycin, oleandomycin, methymycin, neomethymycin, narbomycin, pikromycin, or lankamycin.

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- 12. The method of claim 1 wherein the biologically active compound is an animal feed promotant.
- 13. The method of claim 12 wherein the animal feed promotant is a monensin.

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- 14. The method of claim 12 wherein the monensin is monensin A or monensin B.
  - 15. The method of claim 1 wherein the cell is a prokaryotic cell.

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- 16. The method of claim 15 wherein the prokaryotic cell is a bacterial cell.
- 17. The method of claim 16 wherein the bacterial cell is *Saccharopolyspora*, *Aeromicrobium or Streptomyces*.

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- 18. The method of claim 17 wherein the bacterial cell is a Saccharopolyspora erythraea or an Aeromicrobium erythreum.
- 19. The method of claim 18 wherein the bacterial cell is Streptomyces fradiae,
  25 Streptomyces avermitilis, Streptomyces cinnamonensis, Streptomyces antibioticus,
  Streptomyces venezuelae, Streptomyces violaceoniger, Streptomyces hygroscopicus,
  Streptomyces spp. FR-008, or Streptomyces griseus.
  - 20. The method of claim 1 wherein the cell is a eukaryotic cell.

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21. The method of claim 20 wherein the eukaryotic cell is a plant cell.

- 22. The method of claim 20 wherein the eukaryotic cell is an animal cell.
- 23. The method of claim 22 wherein the animal cell is a mammalian cell.

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- 24. The method of claim 1 wherein inhibiting is accomplished by reducing the level of a co-factor necessary for methylmalonyl-CoA mutase activity.
  - 25. The method of claim 24 wherein the co-factor is coenzyme B12.

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- 26. The method of claim 25 wherein the level of coenzyme B12 is reduced by inhibiting the transcription of a *cob* gene.
- 27. The method of claim 1 wherein inhibiting the activity of methylmalonyl-CoA mutase is accomplished by inhibiting the transcription of a gene for methylmalonyl-CoA mutase.
  - 28. The method of claim 27 wherein inhibiting the transcription of a gene for methylmalonyl-CoA mutase is accomplished by mutating the gene for methylmalonyl-CoA mutase such that the mutated gene does not encode an enzymatically active methylmalonyl-CoA mutase.
  - 29. The method of claim 28 wherein mutating is accomplished by mutating a wild type methylmalonyl CoA gene *in vitro*.

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30. A method of increasing the production of an antibiotic in a bacterial cell wherein the antibiotic is derived at least in part from methylmalonyl-CoA, the method comprising the step of inhibiting the activity of methylmalonyl-CoA mutase in the bacterial cell.

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- 31. The method of claim 30 wherein the antibiotic is a polyketide macrolide antibiotic.
- 32. The method of claim 31 wherein the polyketide macrolide antibiotic is erythromycin.
  - 33. The method of claim 32 wherein the bacterial cell is a *Saccharopolyspora* or *Aeromicrobium*.
- 15 34. The method of claim 33 wherein the bacterial cell is *Saccharopolyspora* erythraea or *Aeromicrobium erythreum*.